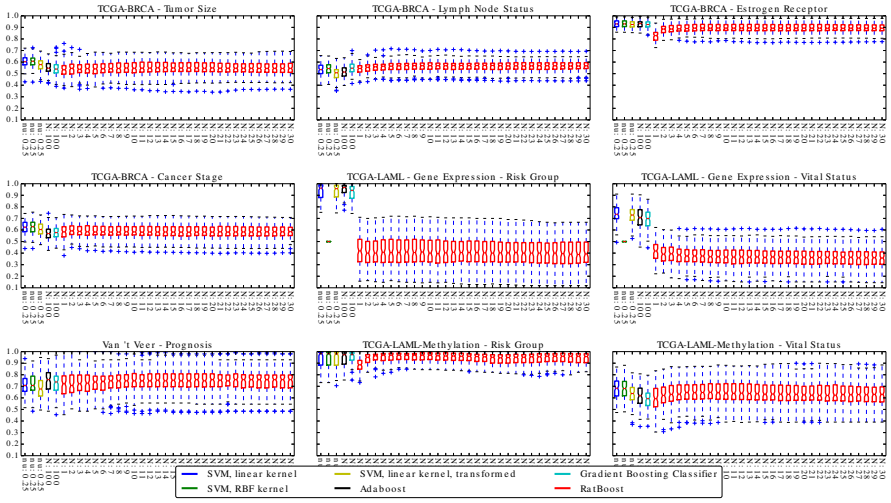
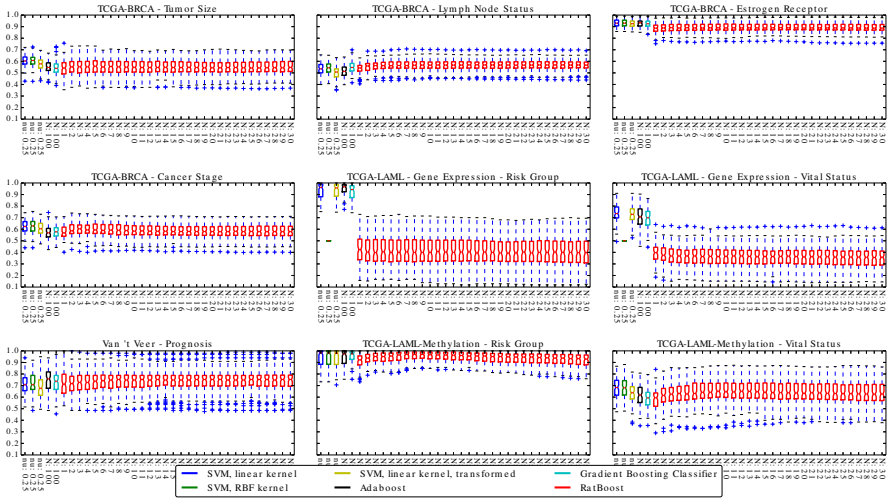
Supplementary Material 1 for: Interpretable per Case Weighted Ensemble Method for Cancer Associations

Adrin Jalali, Nico Pfeifer

1 Method's Performance

The following plots represent the performance of our method with respect to differenct regularization levels, from highly regularized to less regularized.




TCGA-BRCA - Tumor Size
TCGA-BRCA - Lymph Node Status
TCGA-BRCA - Estrogen Receptor
TCGA-BRCA - Cancer Stage
TCGA-LAML - Gene Expression - Risk Group
TCGA-LAML - Gene Expression - Vital Status
Van 't Veer - Prognosis
TCGA-LAML-Methylation - Risk Group
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Gradient Boosting Classifier
SVM, RBF kernel
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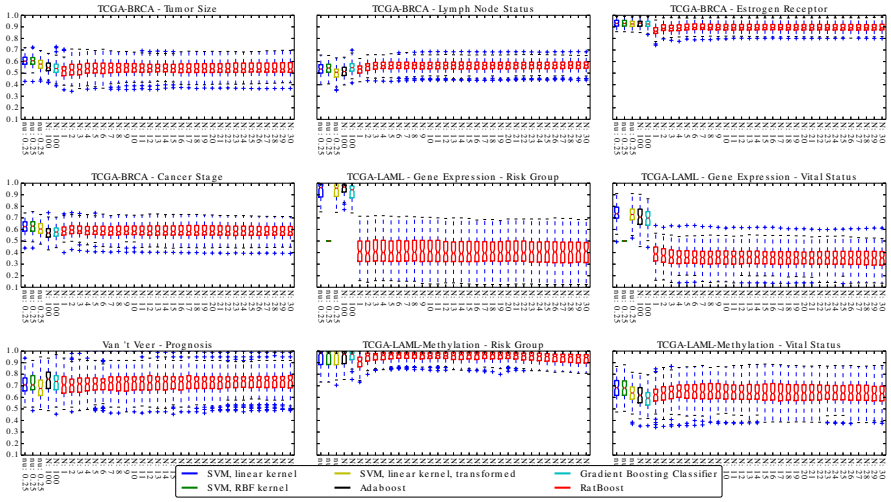
RatBoost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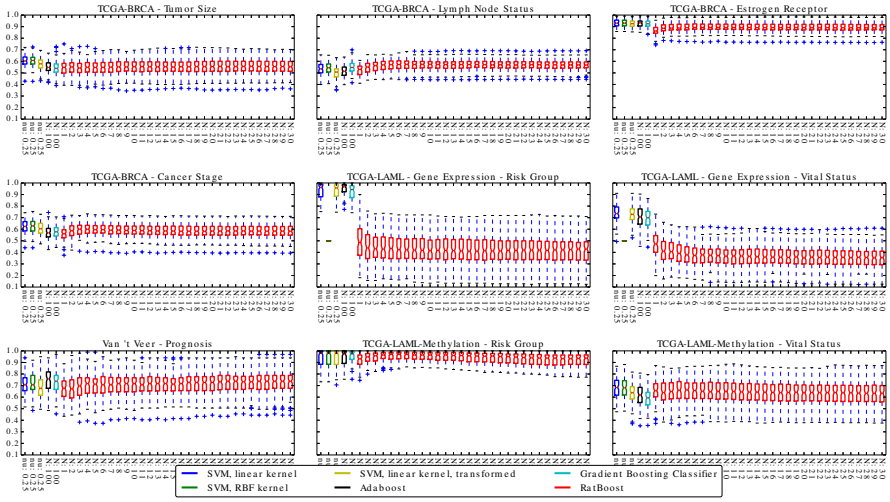
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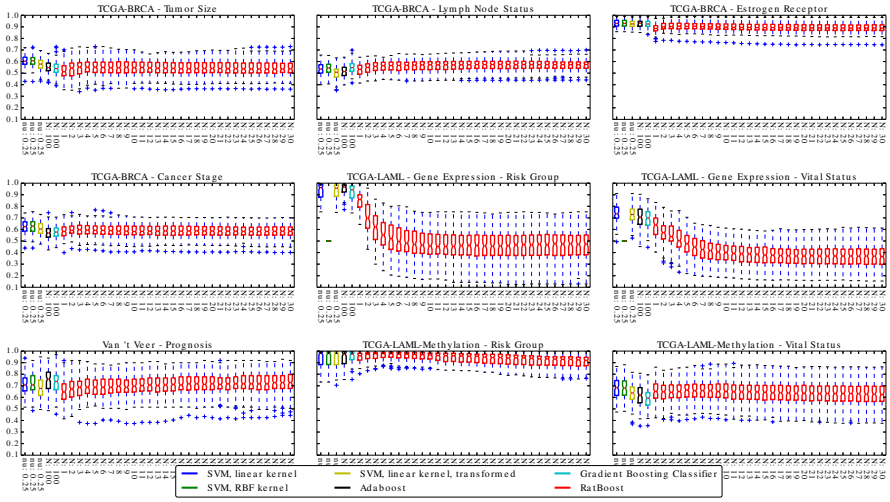
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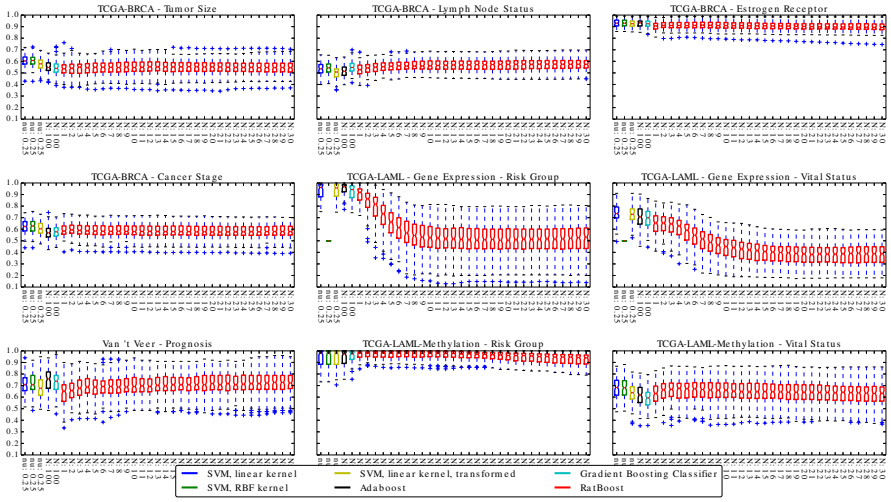
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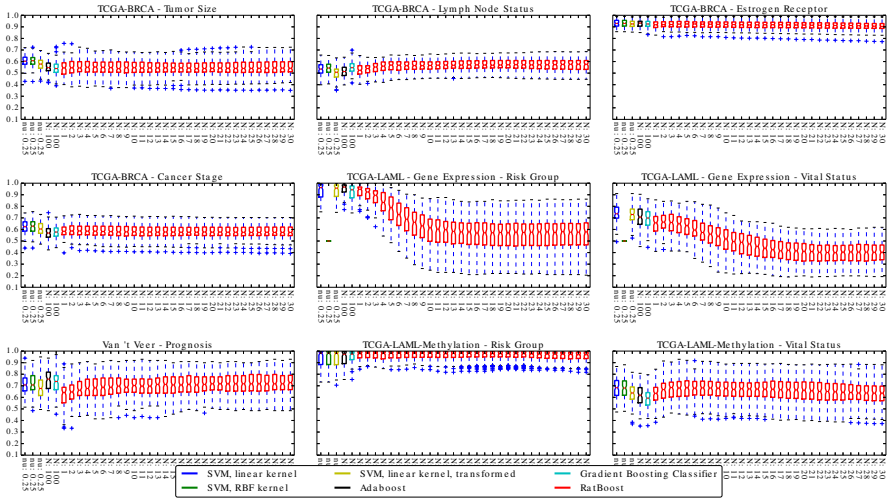
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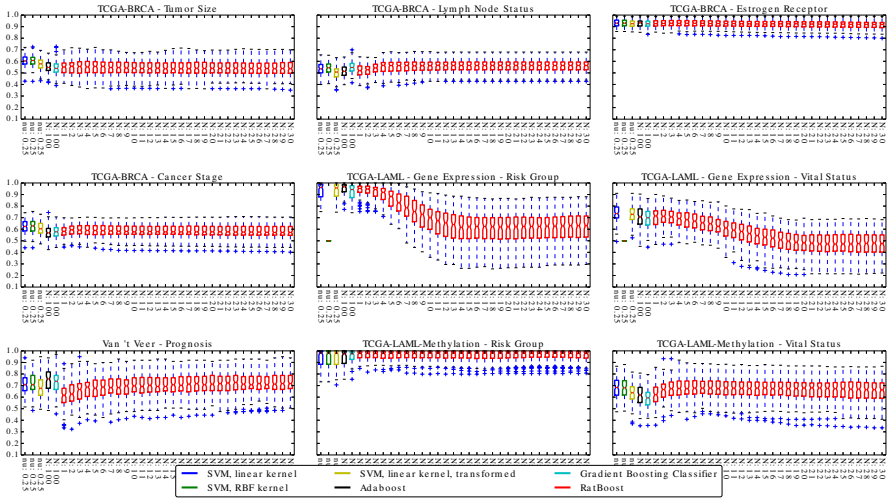
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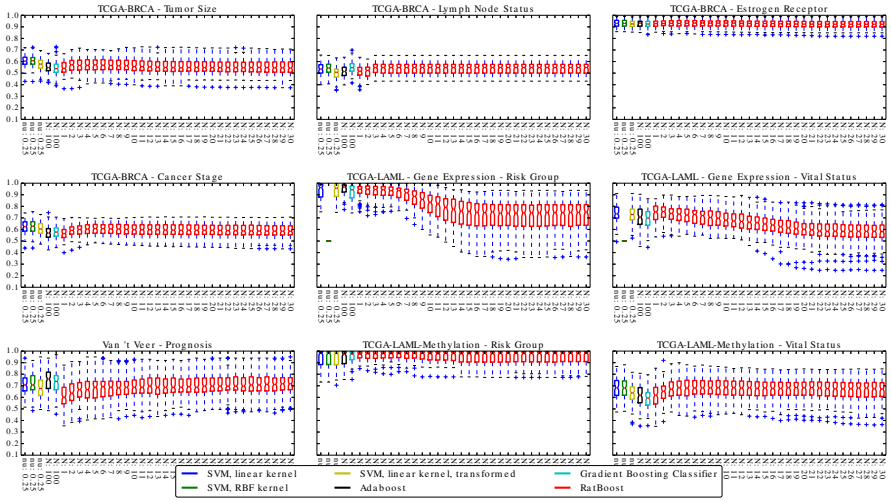
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